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... signal transduction **database**; a knowledge base on signal transduction **networks**

F Schacherer, C Choi, U Gotze, M Krull, S Pistor, ... - ..., 2001 - Oxford Univ Press

... Publications often do not exactly specify which member of a **protein** family was used in an experiment. ... To take into account family and state relationships in the **database**, the **search** can be extended to traverse these relations too. ...

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[BOOK] Protein turnover in mammalian tissues and in the whole body

JC Waterlow, PJ Garlick, DJ Millward, ... - 1978 - cababstractplus.org

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Cited by 839

Kinesin and dynein superfamily **proteins** and the mechanism of organelle transport

N Hirokawa - Science, 1998 - sciencemag.org

... is designated as Cef56E3.3. Thus, it is easy to access **database** entries using ... as retrograde motors in axons, and slow transport motors carrying cytoskeletal **proteins** and cytosolic **proteins**. A systematic **search** for cDNAs encoding as yet unidentified KIFs has been performed ...

Cited by 929 - Related articles - BL Direct - All 8 versions

The geometry of metal-ligand interactions relevant to **proteins**

MM Harding - Acta Crystallographica Section D: Biological ..., 1999 - scripts.iucr.org

... In each **search** of the CSD, the chemical connectivity of a 'query structure' is matched against that of a structure within the **database**. ... It is desirable that the 'small-molecule' structures selected should be good analogues for the metal-ligand interactions in **proteins**, but too ...

Cited by 253 - Related articles - BL Direct - All 11 versions

... of an mRNA encoding the 27 kDa outer dense fiber **protein** of rat spermatozoa

CR Morales, R Oiko, Y Clermont - Molecular Reproduction ..., 1994 - interscience.wiley.com

... remain at high levels from steps 11 to 15, and diminish in steps 16-18 at the time ODF **protein** synthesis and ... A literature and NCBI **database search** revealed that the nucleotide sequence of the 027 cDNA is homologous to the rat gene RT7 ... SEARCH WILEY INTERSCIENCE. ...

Cited by 53 - Related articles - BL Direct - All 3 versions

A pathway finding system for the cell signaling **networks** **database**

T Takai-Igarashi, T Kaminuma - In Silico Biology, 1999 - IOS Press

... The user can **search** molecules by using "CSNDB query" and "Alphabetical lists of molecules". ...[20] Takai-Igarashi, T, Nadaoka, Y, and Kaminuma, T. A **database** for cell ... Guide, Installation Guide, Configuration Guide and Syntactic Definitions for the ACEDB Data Base Manager. ...

Cited by 66 - Related articles - BL Direct - All 5 versions

Detection of dsRNA-binding domains in RNA helicase A and Drosophila maleless: ...

TJ Gibson, JD Thompson - Nucleic acids research, 1994 - Oxford Univ Press

... with single dsRBD sequences perform much less well than the profile **searches**, with many ... Scores and ranking positions resulting from searching the SWISS-PROT **database** with individual ...

RNA-binding domains which, in many cases, will be provided by accessory **proteins** ...

Cited by 80 - Related articles - BL Direct - All 5 versions

**Protein structure prediction: recognition of primary, secondary, and tertiary ...**

F Eisenhaber, B Persson, P ... - Critical reviews in ..., 1995 - informahealthcare.com

... 13 4. Neural Networks .... 14 ... to PROSITEDOC; DE, full name of the pattern; PA; pattern description (x denotes any amino acid); NR, numerical result of SWISS-PROT database search; DR, identifiers of **proteins** containing the ...

Cited by 104 - Related articles - BL Direct - All 7 versions

**Using Bayesian networks to analyze expression data**

N Friedman, M Linial, I Nachman, D ... - Journal of computational ..., 2000 - liebertonline.com

... Examples of other **search** methods that advance using one-arc changes include beam-**search**, stochastic hill ... It remains possible that an unobserved variable (eg, **protein** activation) is an intermediate in their ... of feature is order relations: Is X an ancestor of Y in all the **networks** of a ...

Cited by 1496 - Related articles - BL Direct - All 116 versions

... Partial Characterisation of the cDNA Encoding the Fox Sperm Protein FSA-Acr. 1 ...

S Beaton, J Have, A Cleary, MP ... - Molecular ..., 1995 - interscience.wiley.com

... **Database searches** with the deduced amino acid sequence of cFSA-Acr.1 revealed that the clone has high homology to ... and sequence analysis shows that they may represent alternatively spliced mRNAs coding for other FSA-Acr **proteins** ... **SEARCH WILEY INTERSCIENCE** ...

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